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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/555,587	03/08/2007	Brian E. Jones	GC800-2-US	4393

7590 01/29/2008
Janet K Castaneda
Genencor International Inc
925 Page Mill Road
Palo Alto, CA 94304-1013

EXAMINER

SAIDHA, TEKCHAND

ART UNIT	PAPER NUMBER
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1652

MAIL DATE	DELIVERY MODE
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01/29/2008

PAPER

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.



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Washington, D.C. 20231

SERIAL NUMBER	FILING DATE	FIRST NAMED APPLICANT	ATTORNEY DOCKET NO.

EXAMINER	
ART UNIT	PAPER NUMBER

Please find below a communication from the EXAMINER in charge of this application

This application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 C.F.R. § 1.821(a)(1) and (a)(2). However, this application fails to comply with the requirements of 37 C.F.R. §§ 1.821-1.825 for the reason(s) set forth on the attached Notice To Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures.

APPLICANT IS GIVEN 30 days FROM THE DATE OF THIS LETTER WITHIN WHICH TO COMPLY WITH THE SEQUENCE RULES, 37 C.F.R. §§ 1.821-1.825. Failure to comply with these requirements will result in ABANDONMENT of the application under 37 C.F.R. § 1.821(g). Extensions of time may be obtained by filing a petition accompanied by the extension fee under the provisions of 37 C.F.R. § 1.136. In no case may an applicant extend the period for response beyond the six month statutory period. Direct the response to the undersigned. Applicant is requested to return a copy of the attached Notice to Comply with the response.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Examiner Tekchand Saidha whose telephone number is (703) 305-6595. If the examiner cannot be reached, inquiries can be directed to Supervisory Patent Examiner, Ponnathapu Achutamurthy whose telephone number is (703) 308-3804. The fax number for the organization where this application or proceeding is assigned is (703) 308-4242.

Any inquiry of a general nature or relating to the status of this application should be directed to the Group receptionist whose telephone number is (703) 308-0196.


TEKCHAND SAIDHA
PRIMARY EXAMINER

10/555,587

=====

Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Fri Aug 03 11:25:30 EDT 2007

=====

Reviewer Comments:

<210> 2

<211> 1209

<212> DNA

<213> Unknown

<220>

<223> environmental sample

<221> misc_feature

<222> 734

<223> n = A,T,C or G

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120

ctcggctttt tcgcccattc ggcgctggat gaagaaaatc cagacggccc gggtcataat
180

ttcgcgcttt tagaccaaatt tgctgccctg aaatgggtgc aggaaaatat cgctgctttc
240

ggcggcgacg cggggaatgt cacgctgttt ggcgagtctg ccggggcgcg tagcgtgctt
300

tcgctgctgg cgtcgccgct ggcgaaaaac cttttccaca aaggtattat acaaagcgcc
360

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420

cattacgggc tgcaaaatgc cacagcggat gaactccgcg ctctgcctgc ggatgggctg
480

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 840

The above <222> response only indicates one "n" location (734); however,
 n's are also located at 781 and 784: please explain them.

(from Sequence 3)

<221> VARIANT

<222> 245, 260, 261

<223> Xaa = Any Amino Acid

<400> 3

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			20					25					30		
Thr	Val	Asn	Tyr	Arg	Leu	Gly	His	Leu	Gly	Phe	Phe	Ala	His	Pro	Ala
		35					40						45		
Leu	Asp	Glu	Glu	Asn	Pro	Asp	Gly	Pro	Val	His	Asn	Phe	Ala	Leu	Leu
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65					70				75					80	
Gly	Gly	Asp	Ala	Gly	Asn	Val	Thr	Leu	Phe	Gly	Glu	Ser	Ala	Gly	Ala
			85					90						95	
Arg	Ser	Val	Leu	Ser	Leu	Leu	Ala	Ser	Pro	Leu	Ala	Lys	Asn	Leu	Phe
			100					105					110		
His	Lys	Gly	Ile	Ile	Gln	Ser	Ala	Tyr	Thr	Leu	Pro	Asp	Val	Asp	Arg
		115					120					125			
Lys	Lys	Ala	Leu	Lys	Arg	Gly	Val	Ala	Leu	Ala	Gly	His	Tyr	Gly	Leu
		130				135					140				
Gln	Asn	Ala	Thr	Ala	Asp	Glu	Leu	Arg	Ala	Leu	Pro	Ala	Asp	Gly	Leu

145		150		155		160									
Trp	Ala	Leu	Glu	Gly	Pro	Leu	Asn	Ile	Gly	Pro	Thr	Pro	Ile	Ser	Gly
			165						170					175	
Asp	Val	Val	Leu	Pro	Glu	Pro	Met	Leu	Asp	Ile	Phe	Phe	Ala	Gly	Arg
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				245						250				255	
Asp	Met	Ala	Phe	Xaa	Xaa	Leu	Gly	Phe	Val	Val	Met	Gln	Ala	Gln	Gln
				260						265				270	

The above <222> response is incorrect: while Xaa is located at 245, "Phe" is located at 260 (not Xaa). Xaa's are located at 261 and 262.

Application No: 10555587

Version No: 1.0

Input Set:

Output Set:

Started: 2007-08-01 10:04:38.617

Finished: 2007-08-01 10:04:39.579

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 962 ms

Total Warnings: 3

Total Errors: 8

No. of SeqIDs Defined: 3

Actual SeqID Count: 3

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
E 342	'n' position not defined found at POS: 1926 SEQID(1)
E 342	'n' position not defined found at POS: 1973 SEQID(1)
E 342	'n' position not defined found at POS: 1976 SEQID(1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
E 342	'n' position not defined found at POS: 781 SEQID(2)
E 342	'n' position not defined found at POS: 784 SEQID(2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
E 341	'Xaa' position not defined , SEQID (3) POS (245)
E 341	'Xaa' position not defined SEQID (3) POS (261)
E 341	'Xaa' position not defined SEQID (3) POS (262)

SEQUENCE LISTING

<110> Genencor International, Inc.

Jones, Brian E.

Grant, William D.

Heaphy, Shaun

Rees, Helen C.

Grant, Susan

<120> Novel Lipolytic Enzyme LIP1

<130> GC801-2-PCT

<140> 10555587

<141> 2007-08-01

<150> PCT/US04/014752

<151> 2004-05-12

<150> US 60/469,931

<151> 2003-05-12

<160> 3

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 4313

<212> DNA

<213> Unknown

<220>

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<221> misc_feature

<222> 1926, 1973, 1976

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<213> Unknown

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<221> misc_feature
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<220>
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<221> VARIANT
<222> 245, 260, 261
<223> Xaa = Any Amino Acid

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35 40 45
Leu Asp Glu Glu Asn Pro Asp Gly Pro Val His Asn Phe Ala Leu Leu
50 55 60
Asp Gln Ile Ala Ala Leu Lys Trp Val Gln Glu Asn Ile Ala Ala Phe

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His Lys Gly Ile Ile Gln Ser Ala Tyr Thr Leu Pro Asp Val Asp Arg			
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Gln Asn Ala Thr Ala Asp Glu Leu Arg Ala Leu Pro Ala Asp Gly Leu			
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Trp Ala Leu Glu Gly Pro Leu Asn Ile Gly Pro Thr Pro Ile Ser Gly			
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260	265	270	
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275	280	285	
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340	345	350	
Pro Leu Ser Trp Pro Ala Cys Val Arg Gly Lys Asp Arg Thr Met Arg			
355	360	365	
Leu Gly Val His Ser Arg Ala Arg Phe Lys Val Glu Asn Arg Phe Met			
370	375	380	
Arg Met Arg Met Gln Leu Phe Lys Arg Val Met Lys His His Val Ser			
385	390	395	400
Leu Asp			